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sequence of the polypeptide encoded in the extended ORF in the derived sequence.

Fig. 27 shows the sequence of the HCV cDNA in clone 12f, the segment which overlaps clone 14i, and the amino acids encoded therein.

Fig. 28 shows the sequence of the HCV cDNA in clone 35f, the segment which overlaps clone 39c, and the amino acids encoded therein.

Fig. 29 shows the sequence of the HCV cDNA in clone 19g, the segment which overlaps clone 35f, and the amino acids encoded therein.

Fig. 30 shows the sequence of clone 26g, the segment which overlaps clone 19g, and the amino acids encoded therein.

Fig. 31 shows the sequence of clone 15e, the segment which overlaps clone 26g, and the amino acids encoded therein.

Fig. 32 shows the sequence in a composite cDNA, which was derived by aligning clones 12f through 15e in the 5' to 3' direction; it also shows the amino acids encoded in the continuous ORF.

Fig. 33 shows a photograph of Western blots of a fusion protein, SOD-NANB₅₋₁₋₁, with chimpanzee serum from chimpanzees infected with BB-NANB, HAV, and HBV.

Fig. 34 shows a photograph of Western blots of a fusion protein, SOD-NANB₅₋₁₋₁, with serum from humans infected with NANBV, HAV, HBV, and from control humans.

Fig. 35 is a map showing the significant features of the vector pAB24.

Fig. 36 shows the putative amino acid sequence of the carboxy-terminus of the fusion polypeptide C100-3 and the nucleotide sequence encoding it.

Fig. 37A is a photograph of a coomassie blue stained polyacrylamide gel which identifies C100-3 expressed in yeast.

Fig. 37B shows a Western blot of C100-3 with serum from a NANBV infected human.

Fig. 38 shows an autoradiograph of a Northern blot of RNA isolated from the liver of a BB-NANBV infected chimpanzee, probed with BB-NANBV cDNA of clone 81.

Fig. 39 shows an autoradiograph of NANBV nucleic acid treated with RNase A or DNase I, and probed with BB-NANBV cDNA of clone 81.

Fig. 40 shows an autoradiograph of nucleic acids extracted from NANBV particles captured from infected plasma with anti-NANB₅₋₁₋₁, and probed with ³²P-labeled NANBV cDNA from clone 81.

Fig. 41a and b shows autoradiographs of filters containing isolated NANBV nucleic acids, probed with ³²P-labeled plus and minus strand DNA probes derived from NANBV cDNA in clone 81.

Fig. 41-1 shows the homologies between a polypeptide encoded in HCV cDNA and an NS protein from Dengue flavivirus.

Fig. 43 shows a histogram of the distribution of HCV infection in random samples, as determined by an ELISA screening.

Fig. 44 shows a histogram of the distribution of HCV infection in random samples using two configurations of immunoglobulin-enzyme conjugate in an ELISA assay.

Fig. 45 shows the sequences in a primer mix, derived from a conserved sequence in NS1 of flaviviruses.

Fig. 46 shows the HCV cDNA sequence in clone k9-1, the segment which overlaps the cDNA in Fig. 27, and the amino acids encoded therein.

Fig. 47 shows the sequence in a composite CDNA which was derived by aligning clones k9-1 through 15e in the 5' to 3' direction; it also shows the amino acids encoded in the continuous ORF.

I. Definitions

The term "hepatitis C virus" has been reserved by workers in the field for an heretofore unknown etiologic agent of NANBH. Accordingly, as used herein, "hepatitis C virus" (HCV) refers to an agent causative of NANBH, which agent is a virus characterised by: (i) a positive stranded RNA genome; (ii) said genome comprising an open reading frame (ORF) encoding a polyprotein; and (iii) the portion of said polyprotein corresponding to Figure 14 having at least 40% homology to the amino acid sequence in Figure 14. This agent was formerly referred to as NANBV and/or BB-NANBV. The terms HCV, NANBV, and BB-NANBV are used interchangeably herein, but all refer to the virus as defined above. As an extension of this terminology, the disease caused by HCV, formerly called NANB hepatitis (NANBH), is called hepatitis C. The terms NANBH and hepatitis C may be used interchangeably herein.

The term "HCV", as used herein, denotes a viral species which causes NANBH, and attenuated strains or defective interfering particles derived therefrom. As shown infra., the HCV genome is comprised of RNA. It is known that RNA containing viruses have relatively high rates of spontaneous mutation, i.e., reportedly

FIG. 32-1 COMBINED ORF OF DNAs 12f through 15e

IlePheLysIleArgMetTyrValGlyGlyValGluHisArgLeuGluAlaAlaCysAsn
 1 CCATATTATAAATCAGGATGTACGTGGGAGGGGTCGAACACAGGCTGGAAGCTGCCTGCA
 GGTATAAATTTTAGTCTACATGCACCTCCACAGCTTGTGTCGACCTTCGACGGACGT
 TrpThrArgGlyGluArgCysAspLeuGluAspArgAspArgSerGluLeuSerProLeu
 61 ACTGGACGCGGGGCGAACGTTGCGATCTGGAAGACAGGGACAGGTCCGAGCTCAGCCCGT
 TGACCTGCGCCCGCTTGCAACGCTAGACCTTCTGTCCCTGTCCAGGCTCGAGTGGGCA
 LeuLeuThrThrThrGlnTrpGlnValLeuProCysSerPheThrThrLeuProAlaLeu
 121 TACTGTGACCACTACACAGTGGCAGGTCTCCCGTGTTCCTTCACAACCTACCAGCCT
 ATGACGACTGGTGATGTGTACCGTCCAGGAGGGCACAAGGAAGTGTGGGATGGTGGCA
 SerThrGlyLeuIleHisLeuHisGlnAsnIleValAspValGlnTyrLeuTyrGlyVal
 181 TGTCCACCGGCTCATCCACCTCCACCAGAACATTGTGGACGTGCAGTACTGTACGGGG
 ACAGGTGGCCGGAGTAGGTGGAGGTGGTCTTGTAAACCTGCACGTATGAACATGCCCC
 GlySerSerIleAlaSerTrpAlaIleLysTrpGluTyrValValLeuLeuPheLeuLeu
 241 TGGGGTCAAGCATCGCGTCTGGGCCATTAGTGGGAGTACGTGCTTCCTGTTCCTTC
 ACCCAGTTTCGTAGCGCAGGACCGGTAATTCACCTCATGCAGCAAGAGGACAAGGAAG
 LeuAlaAspAlaArgValCysSerCysLeuTrpMetMetLeuLeuIleSerGlnAlaGlu
 301 TGTTCGACAGCGCGCGTCTGCTCCTGCTTGTGGATGATGCTACTCATATCCCAAGCGG
 ACGAACGTCTGCGCGCGCAGACGAGGACGAACACCTACTACGATGAGTATAGGGTTCGCC
 AlaAlaLeuGluAsnLeuValIleLeuAsnAlaAlaSerLeuAlaGlyThrHisGlyLeu
 361 AGGCGGCTTTGGAGAACCTCGTAATACTTAATGCAGCATCCCTGGCCGGGACGCAAGGTC
 TCCGCCGAAACCTCTTGGAGCATTATGAATTACGTGCTAGGGACCGCCCTGCGTGCCAG
 ValSerPheLeuValPhePheCysPheAlaTrpTyrLeuLysGlyLysTrpValProGly
 421 TTGTATCTTCCTCGTGTCTTCTGCTTTGCATGGTATTGTAAGGGTAAGTGGGTGCCCC
 AACATAGGAAGGAGCACAAGAAGACGAAACGTACCATAAACTTCCATTACCCACGGGC
 AlaValTyrThrPheTyrGlyMetTrpProLeuLeuLeuLeuLeuAlaLeuProGln
 481 GAGCGGTCTACACCTTCTACGGGATGTGGCCTCTCTCTGCTCCTGTTGGCGTTGCCCC
 CTCGCCAGATGTGGAAGATGCCCTACACGGAGAGGAGGACGAGGACAACCGCAACGGGG
 ArgAlaTyrAlaLeuAspThrGluValAlaAlaSerCysGlyGlyValValLeuValGly
 541 AGCGGGCGTACGCGCTGGACACGGAGGTGGCCGCGTCTGTGGCGGTGTTGTTCTCGTCG
 TCGCCCGCATGCGCGACCTGTGCCTCCACCGGCGCAGCACACGCCCACAACAAGAGCAGC
 LeuMetAlaLeuThrLeuSerProTyrTyrLysArgTyrIleSerTrpCysLeuTrpTrp
 601 GGTGTATGGCGCTGACTCTGTCAACCATATTACAAGCGCTATATCAGCTGGTGCTTGTGGT
 CCAACTACCGCGACTGAGACAGTGGTATAATGTTCCGATATAGTCGACCAACGAACCA
 LeuGlnTyrPheLeuThrArgValGluAlaGlnLeuHisValTrpIleProProLeuAsn
 661 GGCTTCAGTATTTTCTGACCAGAGTGAAGCGCAACTGCACGTGTGGATTCCCCCCCTCA
 CCGAAGTCATAAAAGACTGGTCTCACCTTCGCGTTGACGTGCACACCTAAGGGGGGGAGT
 ValArgGlyGlyArgAspAlaValIleLeuLeuMetCysAlaValHisProThrLeuVal
 721 ACGTCCGAGGGGGGCGCGACGCGTCATCTTACTCATGTGTGCTGTACACCGACTCTGG
 TGCAGGCTCCCCCGCGCTGCGGCAGTAGAATGAGTACACACGACATGTGGGCTGAGACC
 PheAspIleThrLysLeuLeuLeuAlaValPheGlyProLeuTrpIleLeuGlnAlaSer
 781 TATTTGACATCACCAATGTCTGCTGGCCGTCTTCGGACCCCTTGGATTCTTCAAGCCA
 ATAACTGTAGTGGTTTAACGACGACCGGCGAGAAGCCTGGGGAAACCTAAGAAGTTCGGT
 LeuLeuLysValProTyrPheValArgValGlnGlyLeuLeuArgPheCysAlaLeuAla
 841 GTTGTCTAAAGTACCCTACTTTGTGCGCGTCCAAGGCCTTCTCCGGTTCTGCGCGTTAG
 CAAACGAATTTTCATGGGATGAAACACGCGCAGGTTCCGGAAGAGGCCAAGACGCGCAATC
 ArgLysMetIleGlyGlyHisTyrValGlnMetValIleIleLysLeuGlyAlaLeuThr
 901 CGCGGAAGATGATCGGAGGCCATTACGTGCAAATGGTCATCATTAAAGTTAGGGGCGCTTA
 GCGCCTTCTACTAGCCTCCGTAATGCACGTTTACCAGTAGTAATTCAATCCCCGCGAAT

961 GlyThrTyrValTyrAsnHisLeuThrProLeuArgAspTrpAlaHisAsnGlyLeuArg
 CTGGCACCTATGTTTATAACCATCTCACTCCTCTTCGGGACTGGGCGCACAAACGGCTTGC
 GACCGTGGATACAAATATTGGTAGAGTGAGGAGAAGCCCTGACCGCGTGTGCGGAACG
 1021 AspLeuAlaValAlaValGluProValValPheSerGlnMetGluThrLysLeuIleThr
 GAGATCTGGCCGTGGCTGTAGAGCCAGTCGTCTTCTCCCAAATGGAGACCAAGCTCATCA
 CTCTAGACCGGCACCGACATCTCGGTACGACAGAGAGGGTTTACCTCTGGTTCGAGTAGT
 1081 TrpGlyAlaAspThrAlaAlaCysGlyAspIleIleAsnGlyLeuProValSerAlaArg
 CGTGGGGGGCAGATACCGCCGCGTGGGTGACATCATCAACGGCTTGCCTGTTTCCGCCC
 GCACCCCCCGTCTATGGCGGCGACGCCACTGTAGTAGTTGCCGAACGGACAAAGGCGGG
 1141 ArgGlyArgGluIleLeuLeuGlyProAlaAspGlyMetValSerLysGlyTrpArgLeu
 GCAGGGGCGGGAGATACTGCTCGGGCCAGCCGATGGAATGGTCTCCAAGGGGTGGAGGT
 CTGCCCCGGCCCTCTATGACGAGCCCGTGGCTACCTTACCAGAGGTTCCCCACCTCCA
 1201 LeuAlaProIleThrAlaTyrAlaGlnGlnThrArgGlyLeuLeuGlyCysIleIleThr
 TGCTGGCGCCCATCACGGCGTACGCCAGCAGACAAGGGGCTCTAGGGTGCTAATCA
 ACGACCGCGGGTAGTGCGCATGCGGGTCTGTTCCCGGAGGATCCACGTATTAGT
 1261 SerLeuThrGlyArgAspLysAsnGlnValGluGlyGluValGlnIleValSerThrAla
 CCAGCCTAACTGGCCGGGACAAAACCAAGTGGAGGGTGAGGTCCAGATTGTGTCAACTG
 GGTCCGATTGACCGGCCCTGTTTGGTTTCACTCCACTCCAGGTCTAACACAGTTGAC
 1321 AlaGlnThrPheLeuAlaThrCysIleAsnGlyValCysTrpThrValTyrHisGlyAla
 CTGCCCCAACCTTCTGGCAACGTGCATCAATGGGGTGTGCTGGACTGTCTACCACGGGG
 GACGGGTTTGGAAGGACCGTTGCACGTAGTTACCCACACGACCTGACAGATGGTGCCCC
 1381 GlyThrArgThrIleAlaSerProLysGlyProValIleGlnMetTyrThrAsnValAsp
 CCGGAACGAGGACCATCGCGTACCCAAGGGTCTGTTCATCCAGATGTATACCAATGTAG
 GGCTTGCTCCTGGTAGCGCAGTGGGTTCCAGGACAGTAGGTCTACATATGGTTACATC
 1441 GlnAspLeuValGlyTrpProAlaProGlnGlySerArgSerLeuThrProCysThrCys
 ACCAAGACCTTGTGGGCTGGCCCGCTCCGCAAGGTAGCCGCTCATTGACACCCTGCACCT
 TGGTTCTGGAACACCGACCGGGCGAGGCGTCCATCGGCGAGTAACTGTGGGACGTGAA
 1501 GlySerSerAspLeuTyrLeuValThrArgHisAlaAspValIleProValArgArgArg
 GCGGCTCCTCGGACCTTTACCTGGTACGAGGCACGCCGATGTCTATCCCGTGGCGCGGC
 CGCGAGGAGCCTGGAAATGGACAGTGCTCCGTGGGCTACAGTAAGGGCACGCGGCGG
 1561 GlyAspSerArgGlySerLeuLeuSerProArgProIleSerTyrLeuLysGlySerSer
 GGGGTGATAGCAGGGGCGAGCTGTGTCGCCCCGGGCCATTCTCTACTGAAAGGCTCCT
 CCCCACATCGTCCCCGTGGACGACAGCGGGGCGGGTAAAGGATGAACCTTCGAGGA
 1621 GlyGlyProLeuLeuCysProAlaGlyHisAlaValGlyIlePheArgAlaAlaValCys
 CGGGGGGTCCGCTGTTGTGCCCCGCGGGGCACGCCGTGGGCATATTTAGGGCGCGGTG
 GCCCCCAGGCGACAACACGGGGCGCCCCGTGCGGCACCGTATAAATCCGGCGCCACA
 1681 ThrArgGlyValAlaLysAlaValAspPheIleProValGluAsnLeuGluThrThrMet
 GCACCCGTGGAGTGGCTAAGGCGGTGGACTTTATCCCTGTGGAGAACCCTAGAGACAACCA
 CGTGGGCACCTCACCGATTCCGCCACCTGAAATAGGGACACCTCTTGGATCTCTGTTGGT
 1741 ArgSerProValPheThrAspAsnSerSerProProValValProGlnSerPheGlnVal
 TGAGGTCCCCGGTGTTCACGGATAACTCCTCTCCACCAAGTAGTCCCCAGAGCTTCCAGG
 ACTCCAGGGGCCACAAGTGCTATTGAGGAGAGGTGGTTCATCAGGGGTCTCGAAGGTCC
 1801 AlaHisLeuHisAlaProThrGlySerGlyLysSerThrLysValProAlaAlaTyrAla
 TGGCTCACTCCATGCTCCACAGGCAGCGCAAAGCACCAAGGTCCCGGCTGCATATG
 ACCGAGTGGAGGTACGAGGGTGTCCGTGCGCGTTTTCGTGGTTCAGGGCGACGTATAC
 1861 AlaGlnGlyTyrLysValLeuValLeuAsnProSerValAlaAlaThrLeuGlyPheGly
 CAGCTCAGGGCTATAAGGTGCTAGTACTCAACCCCTCTGTTGCTGCAACACTGGGCTTGT
 GTCGAGTCCCGATATTCCACGATCATGAGTTGGGGAGACAACGAGCTTGTGACCCGAAAC
 AlaTyrMetSerLysAlaHisGlyIleAspProAsnIleArgThrGlyValArgThrIle

FIG. 32-2

1921 GTGCTTACATGTCCAAGGCTCATGGGATCGATCCTAACATCAGGACCGGGGTGAGAACA
 CACGAATGTACAGGTTCCGAGTACCCTAGCTAGGATTGTAGTCCTGGCCCCACTCTTGTT
 ThrThrGlySerProIleThrTyrSerThrTyrGlyLysPheLeuAlaAspGlyGlyCys
 1981 TTACCACTGGCAGCCCCATCACGTACTCCACCTACGGCAAGTTCCTTGCCGACGGCGGGT
 AATGGTGACCGTCGGGGTAGTGCATGAGGTGGATGCCGTTCAAGGAACGGCTGCGCCCA
 SerGlyGlyAlaTyrAspIleIleIleCysAspGluCysHisSerThrAspAlaThrSer
 2041 GCTCGGGGGGCGCTTATGACATAAATTTGTGACGAGTGCCACTCCACGGATGCCACAT
 CGAGCCCCCGGAATACTGTATTATTAAACACTGCTACGGTGAGGTGCTACGGTGTA
 IleLeuGlyIleGlyThrValLeuAspGlnAlaGluThrAlaGlyAlaArgLeuValVal
 2101 CCATCTTGGGCATCGGCCTGTCTTGACCAAGCAGAGACTGCGGGGGGAGACTGGTTG
 GGTAGAACCCGTAGCCGTGACAGGAAGTGGTTGCTCTGACGCCCCGCTCTGACCAAC
 LeuAlaThrAlaThrProProGlySerValThrValProHisProAsnIleGluGluVal
 2161 TGCTCGCCACCGCCACCCCTCCGGGCTCCGTCACTGTGCCCCATCCCAACATCGAGGAGG
 ACGAGCGGTGGCGGTGGGGAGGCCGAGGCAGTGACACGGGGTAGGGTTGTAGCTCCTCC
 AlaLeuSerThrThrGlyGluIleProPheTyrGlyLysAlaIleProLeuGluValIle
 2221 TTGCTCTGTCCACCACCGGAGAGATCCCTTTTTACGGCAAGGCTATCCCCCTCGAAGTAA
 AACGAGACAGGTGGTGGCCTCTCTAGGGAAAAATGCCGTTCCGATAGGGGGAGCTTCATT
 LysGlyGlyArgHisLeuIlePheCysHisSerLysLysLysCysAspGluLeuAlaAla
 2281 TCAAGGGGGGAGACATCTCATCTTCTGTCAATTCAAAGAAGAAGTGCGACGAACTCGCG
 AGTTCCCCCCTCTGTAGAGTAGAAGACAGTAAGTTCTTCTTCACGCTGCTTGAGCGGC
 LysLeuValAlaLeuGlyIleAsnAlaValAlaTyrTyrArgGlyLeuAspValSerVal
 2341 CAAAGCTGGTGCATTGGGCATCAATGCCGTGGCCTACTACCGCGTCTTGACGTGTCCG
 GTTTCGACCAGCGTAACCCGTAGTTACGGCACCGGATGATGGCGCCAGAACTGCACAGGC
 IleProThrSerGlyAspValValValAlaThrAspAlaLeuMetThrGlyTyrThr
 2401 TCATCCCGACCAGCGGGATGTTGTGCTGTGGCAACCGATGCCCTCATGACCGGTATA
 AGTAGGGCTGGTGGCGCTACAACAGCAGCACCGTTGGCTACGGGAGTACTGGCGGATAT
 GlyAspPheAspSerValIleAspCysAsnThrCysValThrGlnThrValAspPheSer
 2461 CCGGGGACTTCGACTCGGTGATAGACTGCAATACGTGTGTACCCAGACAGTCGATTTC
 GCGCGCTGAAGCTGAGCCACTATCTGACGTTATGCACACAGTGGGTCTGTGCTAAAGT
 LeuAspProThrPheThrIleGluThrIleThrLeuProGlnAspAlaValSerArgThr
 2521 GCCTTGACCTACCTTCACCATGAGACAATCAGCTCCCCCAGGATGCTGTCTCCGCA
 CGGAACCTGGGATGGAAGTGGTAACCTCTGTAGTGCGAGGGGGTCTACGACAGAGGGGT
 GlnArgArgGlyArgThrGlyArgGlyLysProGlyIleTyrArgPheValAlaProGly
 2581 CTCAACGTCGGGGCAGGACTGGCAGGGGGAAGCCAGGCATCTACAGATTGTGGCACCGG
 GAGTTGCAGCCCGTCTGACCGTCCCCCTCGGTCCGTAGATGTCTAAACACCGTGGCC
 GluArgProSerGlyMetPheAspSerSerValLeuCysGluCysTyrAspAlaGlyCys
 2641 GGGAGCGCCCTCCGGCATGTTGACTCGTCCGTCCTCTGTGAGTGCTATGACGAGGCT
 CCTCGCGGGGAGGCGGTACAAGCTGAGCAGGCAGGAGACACTACGATACTGCGTCCGA
 AlaTrpTyrGluLeuThrProAlaGluThrThrValArgLeuArgAlaTyrMetAsnThr
 2701 GTGCTTGGTATGAGCTCACGCCCGCGAGACTACAGTTAGGCTACGAGCGTACATGAACA
 CACGAACCATACTCGAGTGCGGGCGGCTCTGATGTCAATCOGATGCTGCGATGTACTTGT
 ProGlyLeuProValCysGlnAspHisLeuGluPheTrpGluGlyValPheThrGlyLeu
 2761 CCCCGGGGCTTCCCGTGTGCCAGGACCATCTTGAATTTGGGAGGGGCTTTTACAGGCC
 GGGGCCCCGAAGGGCACACGGTCTCTGGTAGAAGTTAAACCCCTCCCGCAGAAATGTCGG
 ThrHisIleAspAlaHisPheLeuSerGlnThrLysGlnSerGlyGluAsnLeuProTyr
 2821 TCACTCATATAGATGCCACTTCTATCCAGACAAAGCAGAGTGGGAGAACCTTCTCT
 AGTGAGTATATCTACGGGTGAAGATAGGGTCTGTTTCTGCTCACCCCTCTTGAAGGAA
 LeuValAlaTyrGlnAlaThrValCysAlaArgAlaGlnAlaProProProSerTrpAsp
 2881 ACCTGGTAGCGTACCAAGCCACCGTGTGCGCTAGGGCTCAAGCCCCCTCCCCATCGTGGG
 TGGACCATCGCATGGTTGCGGTGGCACACGCGATCCGAGTTCGGGGAGGGGGTAGCACCC

FIG. 32-3

2941 GlnMetTrpLysCysLeuIleArgLeuLysProThrLeuHisGlyProThrProLeuLeu
 ACCAGATGTGGAAGTGTGTTGATTGCGCTCAAGCCCACCCTCCATGGGCCAACCCCCCTGC
 TGGTCTACACCTTCACAACTAAGCGGAGTTCGGGTGGGAGGTACCCGGTTGTGGGGACG

3001 TyrArgLeuGlyAlaValGlnAsnGluIleThrLeuThrHisProValThrLysTyrIle
 TATACAGACTGGGCGCTGTTTCAAGTAAATCACCTGACGCACCCAGTCACCAAATACA
 ATATGTCTGACCGCGACAAGTCTTACTTTAGTGGGACTGCGTGGGTTCAGTGGTTTATGT

3061 MetThrCysMetSerAlaAspLeuGluValValThrSerThrTrpValLeuValGlyGly
 TCATGACATGCATGTGGGCCGACCTGGAGGTGCTCACGAGCACCTGGGTGCTCGTTGGCG
 AGTACTGTACGTACAGCCGGCTGGACCTCCAGCAGTGTCTGTTGGACCCAGAGCAACCGC

3121 ValLeuAlaAlaLeuAlaAlaTyrCysLeuSerThrGlyCysValValIleValGlyArg
 GCGTCTGGCTGCTTTGGCGCGTATTGCGTGTCAACAGGCTGCGTGGTCATAGTGGGCA
 CGCAGGACCGACGAAACCGGCGCATAACGGACAGTTGTCCGACGCACCACTATCACCCGT

3181 ValValLeuSerGlyLysProAlaIleIleProAspArgGluValLeuTyrArgGluPhe
 GGGTGTCTTGTTCGGAAGCCGGCAATCATACCTGACAGGAAGTCTCTACCGAGAGT
 CCCAGCAGAACAGGCCCTTGGCGGTTAGTATGGACTGTCCCTTCAGGAGATGGCTCTCA

3241 AspGluMetGluGluCysSerGlnHisLeuProTyrIleGluGlnGlyMetMetLeuAla
 TCGATGAGATGGAAGAGTGTCTCAGCACTTACCGTACATCGAGCAAGGGATGATGCTCG
 AGCTACTCTACCTTCTCAGGAGTGTGTAATGGCATGTAGTCTGTTCCCTACTACGAGC

3301 GluGlnPheLysGlnLysAlaLeuGlyLeuLeuGlnThrAlaSerArgGlnAlaGluVal
 CCGAGCAGTTCAAGCAGAAGGCCCTCGGCCCTCTGCAGACCGCGTCCCGTCAGGCAGAGG
 GGCTCGTCAAGTTCGTCTTCGGGGAGCCGGAGGACGTCTGGCGCAGGGCAGTCCGTCTCC

3361 IleAlaProAlaValGlnThrAsnTrpGlnLysLeuGluThrPheTrpAlaLysHisMet
 TTATCGCCCCCTGTGTCCAGACCACTGGCAAACTCGAGACCTTCTGGGCGAAGCATA
 AATAGCGGGGACGACAGGTCTGGTTGACCGTTTTTGTAGCTCTGGAAGACCCGCTTGTAT

3421 TrpAsnPheIleSerGlyIleGlnTyrLeuAlaGlyLeuSerThrLeuProGlyAsnPro
 TGTGGAACCTTCATCAGTGGGATACAATACTTGGCGGGCTGTGTAACGCTGCCTGGTAACC
 ACACCTTGAAGTAGTCACCTTATGTTATGAACGCCCGAACAGTTGCGACGGACCATGG

3481 AlaIleAlaSerLeuMetAlaPheThrAlaAlaValThrSerProLeuThrThrSerGln
 CCGCCATTGCTTCATTGATGGCTTTTACAGCTGCTGTACCAGCCCACTAACCACTAGCC
 GCGGTAACGAAGTAACCTACCGAAAATGTGACGACAGTGGTGGGTGATTGGTGATCGG

3541 ThrLeuLeuPheAsnIleLeuGlyGlyTrpValAlaAlaGlnLeuAlaAlaProGlyAla
 AAACCTCTCTTCAACATATTGGGGGGGTGGGTGGCTGCCAGCTCGCGGCCCCGGGTG
 TTTGGGAGGAGAAGTTGTATAACCCCCCAACCCACCGACGGGTGAGCGGGGGGGCCAC

3601 AlaThrAlaPheValGlyAlaGlyLeuAlaGlyAlaAlaIleGlySerValGlyLeuGly
 CCGCTACTGCCTTTGTGGGCGCTGGCTTAGCTGGCGCGCCATCGGCAGTGTGGACTGG
 GGCGATGACGGAAACACCGCGACCGAATCGACCGGGCGGTAGCCGTCAACCTGACC

3661 LysValLeuIleAspIleLeuAlaGlyTyrGlyAlaGlyValAlaGlyAlaLeuValAla
 GGAAGTCTCATAGACATCTTGCAGGTATGGCGGGGGGTGGCGGGAGCTCTGTGG
 CCTTCCAGGAGTATCTGTAGGAACGTCCCATACCGCGCCCGCACCGCCCTCGAGAACACC

3721 PheLysIleMetSerGlyGluValProSerThrGluAspLeuValAsnLeuLeuProAla
 CATTCAAGATCATGAGCGGTGAGGTCCCTCCACGGAGGACCTGGTCAATCTACTGCCCG
 GTAAGTCTAGTACTCGCCACTCCAGGGGAGGTGCTCTGGACCAGTTAGATGACGGGG

3781 IleLeuSerProGlyAlaLeuValValGlyValValCysAlaAlaIleLeuArgArgHis
 CCATCTCTCGCCCGGAGCCCTGTAGTCGGCGTGGTCTGTGCAGCAATACTGCGCGCGC
 GGTAGGAGAGCGGGCCTCGGGAGCATCAGCGCACCAAGACAGTCTTATGACGCGGGCG

3841 ValGlyProGlyGluGlyAlaValGlnTrpMetAsnArgLeuIleAlaPheAlaSerArg
 ACGTTGGCCCGGCGAGGGGGCAGTGCAGTGAACCGGCTGATAGCCTTCGCCTCCC
 TGCAACCGGGCCCGCTCCCCCGTCACGTACCTACTTGGCCGACTATCGGAAGCGGAGGG

GlyAsnHisValSerProThrHisTyrValProGluSerAspAlaAlaAlaArgValThr

FIG. 32-4

3901 GGGGGAACCATGTTTCCCCACGCACTACGTGCCGGAGAGCGATGCAGCTGCCCCGCGTCA
 CCCCCTTGGTACAAAGGGGGTGCGTGATGCACGGCCTCTCGCTACGTGACGGGCGCAGT
 AlaIleLeuSerSerLeuThrValThrGlnLeuLeuArgArgLeuHisGlnTrpIleSer
 3961 CTGCCATACTCAGCAGCCTCACTGTAACCCAGCTCCTGAGGCGACTGCACCAGTGGATAA
 GACGGTATGAGTCGTGGAGTGACATTGGGTCGAGGACTCCGCTGACGTGGTCACCTATT
 SerGluCysThrThrProCysSerGlySerTrpLeuArgAspIleTrpAspTrpIleCys
 4021 GCTCGGAGTGTAACCACTCCATGCTCGGTTCTGGCTAAGGGACATCTGGGACTGGATAT
 CGAGCCTCACATGGTGAGGTACGAGGCCAAGGACCGATTCCCTGTAGACCCTGACCTATA
 GluValLeuSerAspPheLysThrTrpLeuLysAlaLysLeuMetProGlnLeuProGly
 4081 GCGAGGTGTTGAGCGACTTTAAGACCTGGCTAAAAGCTAAGCTCATGCCACAGCTGCCTG
 CGCTCCACAACCTCGCTGAAATTCTGGACCGATTTCGATTGAGTACGGTGTGACGGAC
 IleProPheValSerCysGlnArgGlyTyrLysGlyValTrpArgValAspGlyIleMet
 4141 GGATCCCCTTGTGTCTCTGCCAGCGCGGGTATAAGGGGGTCTGGCGAGTGGACGGCATCA
 CCTAGGGGAAACACAGGACGGTGGCGCCATATTCCCCAGACCGCTCACCTGCCGTAGT
 HisThrArgCysHisCysGlyAlaGluIleThrGlyHisValLysAsnGlyThrMetArg
 4201 TGCACACTCGCTGCCACTGTGGAGCTGAGATCACTGGACATGTCAAAAACGGGACGATGA
 ACGTGTGAGCGACGGTGACACCTCGACTCTAGTGACCTGTACAGTTTTGGCCCTGCTACT
 IleValGlyProArgThrCysArgAsnMetTrpSerGlyThrPheProIleAsnAlaTyr
 4261 GGATCGTCGGTCTTAGGACCTGCAGGAACATGTGGAGTGGGACCTTCCCCATTAAATGCCT
 CCTAGCAGCCAGGATCCTGGACGTCTTGTACACCTCACCTGGAAGGGGTAATTACGGA
 ThrThrGlyProCysThrProLeuProAlaProAsnTyrThrPheAlaLeuTrpArgVal
 4321 ACACCACGGGCCCTGTACCCCTCTCTGCGCGAAGTACACGTTGCGGCTATGGAGGG
 TGTGGTGCCCGGGACATGGGGGAAGGACGGGCTTGATGTGAAGCGGATACCTCCC
 SerAlaGluGluTyrValGluIleArgGlnValGlyAspPheHisTyrValThrGlyMet
 4381 TGTCTGCAGAGGAATATGTGGAGATAAGGCAGGTGGGGGACTTCCACTACGTGACGGGTA
 ACAGACGTCTCCTTATACACCTCTATTCCGTCCACCCCTGAAGGTGATGCACTGCCCAT
 ThrThrAspAsnLeuLysCysProCysGlnValProSerProGluPhePheThrGluLeu
 4441 TGACTACTGACAATCTCAAATGCCGTGCCAGGTCCCATCGCCCGAATTTTTCACAGAAT
 ACTGATGACTGTAGAGTTTACGGGCACGGTCCAGGGTAGCGGGCTAAAAAGTGCTTA
 AspGlyValArgLeuHisArgPheAlaProProCysLysProLeuLeuArgGluGluVal
 4501 TGGACGGGGTGCGCCTACATAGGTTTGCGCCCCCTGCAAGCCCTGCTGCGGGAGGAGG
 ACCTGCCCAOCCGGATGTATCCAAACGCGGGGGGACGTTCCGGGAACGAGGCCCTCTCC
 SerPheArgValGlyLeuHisGluTyrProValGlySerGlnLeuProCysGluProGlu
 4561 TATCATTCAGAGTAGGACTCCACGAATACCGGTAGGGTCCGAATTACCTTGCAGAGCCG
 ATAGTAAGTCTCATCTGAGGTGCTTATGGGCCATCCAGCGTTAATGGAACGCTCGGGC
 ProAspValAlaValLeuThrSerMetLeuThrAspProSerHisIleThrAlaGluAla
 4621 AACCGGACGTGGCCGTGTGACGTCCATGCTCACTGATCCCTCCCATATAACAGCAGAGG
 TTGGCCTGCACCGGCACAACCTGCAGGTACGAGTGACTAGGGAGGGTATATTGTCGTCTCC
 AlaGlyArgArgLeuAlaArgGlySerProProSerValAlaSerSerSerAlaSerGln
 4681 CGGCCGGGCGAAGGTGGGAGGGGATCACCCCTCTGTGGCCAGCTCCTCGGCTAGCC
 GCGGGCCCGCTTCCAACCGCTCCCTAGTGGGGGAGACACCGGTGAGGAGCCGATCGG
 LeuSerAlaProSerLeuLysAlaThrCysThrAlaAsnHisAspSerProAspAlaGlu
 4741 AGCTATCCGCTCCATCTCTCAAGGCAACTTGACCGCTAACCATGACTCCCTGATGCTG
 TCGATAGGCGAGGTAGAGAGTTCCGTGAACGTGGCGATTGGTACTGAGGGGACTACGAC
 LeuIleGluAlaAsnLeuLeuTrpArgGlnGluMetGlyGlyAsnIleThrArgValGlu
 4801 AGCTCATAGAGCCAACCTCCTATGGAGGCAGAGATGGGCGGCAACATCACCAGGGTGG
 TCGATATCTCCGTTGGAGGATACCTCCGTCTCTACCGCGGTTGTAGTGGTCCCAAC
 SerGluAsnLysValValIleLeuAspSerPheAspProLeuValAlaGluGluAspGlu
 4861 AGTCAGAAAACAAAGTGGTGATTCTGGACTCCTTCGATCCGCTTGTGGCGGAGGAGGACG
 TCAGTCTTTTTCACCACCTAAGACCTGAGGAAGCTAGGCGAACACCGCCTCTCTCTGC

FIG. 32-5

4921 ArgGluIleSerValProAlaGluIleLeuArgLysSerArgArgPheAlaGlnAlaLeu
 AGCGGGAGATCTCCGTACCCGAGAAATCTTGCGGAAGTCTCGGAGATTGCCCCAGGCCC
 TCGCCCTCTAGAGGCATGGGCGTCTTTAGGACGCCCTTCAGAGCCTCTAAGCGGGTCCGGG

 4981 ProValTrpAlaArgProAspTyrAsnProProLeuValGluThrTrpLysLysProAsp
 TGCCCGTTTGGGCGGGCCGGACTATAACCCCCCGCTAGTGGAGACGTGGAAAAAGCCCG
 ACGGGCAAACCCGCGCGGCCCTGATATTGGGGGGCGATCACCTCTGCACCTTTTTCGGGGC

 5041 TyrGluProProValValHisGlyCysProLeuProProProLysSerProProValPro
 ACTACGAACCACCTGTGGTCCATGGCTGTCCGCTTCCACCTCCAAAGTCCCTCTCTGTGC
 TGATGCTTGGTGGACACCAGGTACCGACAGGCGAAGGTGGAGGTTTCAGGGGAGGACACG

 5101 ProProArgLysLysArgThrValValLeuThrGluSerThrLeuSerThrAlaLeuAla
 CTCGCCCTCGGAAGAAGCGGACGGTGGTCTCACTGAATCAACCCTATCTACTGCCTTGG
 GAGGCGGAGCCTTCTTCGCTGCCACCAGGAGTGAAGTTAGTTGGGATAGATGACGGAACC

 5161 GluLeuAlaThrArgSerPheGlySerSerSerThrSerGlyIleThrGlyAspAsnThr
 CCGAGCTCGCCACCAGAAGCTTTGGCAGCTCCTCAACTTCGGGCATTACGGGCGACAATA
 GGCTCGAGCGGTGGTCTTCGAAACCGTCGAGGAGTTGAAGGCCGTAATGCCGCTGTTAT

 5221 ThrThrSerSerGluProAlaProSerGlyCysProProAspSerAspAlaGluSerTyr
 CGACAACATCCTCTGAGCCCGCCCTTCTGGCTGCCCCCGGACTCCGACGCTGAGTCCT
 GCTGTGTAGGAGACTCGGGCGGGGAAGACCACGGGGGGGCTGAGGCTGCGACTCAGGA

 5281 SerSerMetProProLeuGluGlyGluProGlyAspProAspLeuSerAspGlySerTrp
 ATTCTCCATGCCCCCTGGAGGGGGAGCCTGGGGATCCGGATCTTAGCGACGGGTGAT
 TAAGGAGGTACGGGGGGACCTCCCCCTCGGACCCCTAGGCCTAGAATCGCTGCCCAGTA

 5341 SerThrValSerSerGluAlaAsnAlaGluAspValValCysCysSerMetSerTyrSer
 GGTCAACGGTCAGTAGTGAGGCCAACGCGGAGGATGTCGTGTGCTGCTCAATGTCTTACT
 CCAGTTGCCAGTCATCACTCCGGTTGCGCCTCCTACAGCACACGACGAGTTACAGAATGA

 5401 TrpThrGlyAlaLeuValThrProCysAlaAlaGluGluGlnLysLeuProIleAsnAla
 CTGGACAGGGCACTCGTCACCCCGTGGCGCGGAAGAACAGAACTGCCCATCAATG
 GAACCTGTCCGGTGAGCAGTGGGGCACGGCGCCTTCTTGTCTTTGACGGGTAGTTAC

 5461 LeuSerAsnSerLeuLeuArgHisHisAsnLeuValTyrSerThrThrSerArgSerAla
 CACTAAGCAACTCGTTGCTACGTACCCACAATTGGGTGATTCCACCACCTCACGCAGTG
 GTGATTGCTTGAGCAACGATGCAGTGGTGTAAACCACATAAGGTGGTGGAGTGCCTCAC

 5521 CysGlnArgGlnLysLysValThrPheAspArgLeuGlnValLeuAspSerHisTyrGln
 CTTGCCAAAGGCAGAAGAAAGTCAATTTGACAGACTGCAAGTTCTGGACAGCCATTACC
 GAACGGTTTTCGTCTTCTTTCAAGTGTAACTGTCTGACGTTCAAGACCTGTGCGGTAATGG

 5581 AspValLeuLysGluValLysAlaAlaAlaSerLysValLysAlaAsnLeuLeuSerVal
 AGGACGTACTCAAGGAGGTTAAAGCAGCGGCGTCAAAAGTGAAGGCTAACTTGCTATCCG
 TCCTGCATGAGTTCTCAATTTCTGTCGCGCAGTTTTCACTTCCGATTGAACGATAGGC

 5641 GluGluAlaCysSerLeuThrProProHisSerAlaLysSerLysPheGlyTyrGlyAla
 TAGAGGAAGCTTGACGCTGACGCCCCACACTCAGCCAAATCCAAGTTTGGTTATGGGG
 ATCTCCTTCGAACGTCGACTGCGGGGTGTGAGTCGGTTTAGGTTCAAACCAATACCCC

 5701 LysAspValArgCysHisAlaArgLysAlaValThrHisIleAsnSerValTrpLysAsp
 CAAAAGACGTCCGTTGCCATGCCAGAAAGGCCGTAACCCACATCAACTCCGTGTGGAAG
 GTTTTCTGCAGGCAACGGTACGGTCTTTCCGGCATGGGTGTAGTTGAGGCACACCTTTC

 5761 LeuLeuGluAspAsnValThrProIleAspThrThrIleMetAlaLysAsnGluValPhe
 ACCTTCTGGAAGACAATGTAACACCAATAGACACTACCATCATGGCTAAGAACGAGGTTT
 TGAAGACCTTCTGTTACATTGTGGTTATCTGTGATGGTAGTACCGATTCTTGCTCCAA

 5821 CysValGlnProGluLysGlyGlyArgLysProAlaArgLeuIleValPheProAspLeu
 TCTGCGTTTACGCTGAGAAGGGGGGTGTAAGCCAGCTCGTCTCATCGTGTTCCTCGATC
 AGACGCAAGTCGACTCTTCCCCCAGCATTCGGTCGAGCAGAGTAGCACAAGGGGCTAG

 GlyValArgValCysGluLysMetAlaLeuTyrAspValValThrLysLeuProLeuAla

FIG. 32-6

5881 TGGGCGTGC GCGTGTGCGAAAAGATGGCTTTGTACGACGTGGTTACAAAGCTCCCCCTGG
 ACCGCGACGCGCACACGCTTTTCTACCGAAACATGCTGCACCAATGTTTCGAGGGGAACC

 ValMetGlySerSerTyrGlyPheGlnTyrSerProGlyGlnArgValGluPheLeuVal
 5941 CCGTGATGGGAAGCTCCTACGGATTCCAATACTCACCAGGACAGCGGGTTGAATTCCTCG
 GGCCTACCTTCGAGGATGCCTAAGGTTATGAGTGGTCTGTGCGCCAACTTAAGGAGC

 GlnAlaTrpLysSerLysLysThrProMetGlyPheSerTyrAspThrArgCysPheAsp
 6001 TGCAAGCGTGGAAGTCCAAGAAAACCCCAATGGGGTTCTCGTATGATACCGCTGCTTTG
 ACGTTCCGACCTTCAGGTTCTTTTGGGGTTACCCCAAGAGCATACTATGGGCGACGAAAC

 SerThrValThrGluSerAspIleArgThrGluGluAlaIleTyrGlnCysCysAspLeu
 6061 ACTCCACAGTCACTGAGAGCGACATCCGTACGAGGAGGCAATCTACCAATGTTGTGACC
 TGAGGTGTCAGTGA CTCTCGCTGTAGGCATGCCCTCCTCGTTAGATGGTTACAACACTGG

 AspProGlnAlaArgValAlaIleLysSerLeuThrGluArgLeuTyrValGlyGlyPro
 6121 TCGACCCCAAGCCCGCGTGGCCATCAAGTCCCTCACCAGAGAGGCTTTATGTTGGGGGCC
 AGCTGGGGGTTCCGGCGCACGGTAGTTTCAGGGAGTGGCTCTCGAAATACAACCCCGG

 LeuThrAsnSerArgGlyGluAsnCysGlyTyrArgArgCysArgAlaSerGlyValLeu
 6181 CTCTTACCAATTCAAGGGGGGAGAACTGCGGCTATCGCAGGTGCGCGCGAGCGGCGTAC
 GAGAATGGTTAAGTTCCCCCTCTTGACGCCGATAGCGTCCACGGCGCGCTCGCCGATG

 ThrThrSerCysGlyAsnThrLeuThrCysTyrIleLysAlaArgAlaAlaCysArgAla
 6241 TGACAACTAGCTGTGGTAACACCCTCACTTGCTACATCAAGGCCGGGCGAGCCTGTGAG
 ACTGTTGATCGACACCATTGTGGGAGTGAACGATGTAGTTCCGGGCCCGTGGACAGCTC

 AlaGlyLeuGlnAspCysThrMetLeuValCysGlyAspAspLeuValValIleCysGlu
 6301 CCGCAGGGCTCCAGGACTGCACCATGCTCGTGTGTGGCGACGACTTAGTCGTTATCTGTG
 GCGTCCCGAGGTCTGACGTGGTACGAGCACACACCGCTGCTGAATCAGCAATAGACAC

 SerAlaGlyValGlnGluAspAlaAlaSerLeuArgAlaPheThrGluAlaMetThrArg
 6361 AAAGCGCGGGGGTCCAGGAGGACGCGCGAGCCTGAGAGCCTTCACGGAGGCTATGACCA
 TTTCCGCGCCCCAGGTCTCTGCGCGCTCGGACTCTCGGAAGTGCTCCGATACTGGT

 TyrSerAlaProProGlyAspProProGlnProGluTyrAspLeuGluLeuIleThrSer
 6421 GGTACTCCGCCCCCTGGGGACCCCCACAACCAATACGACTTGGAGCTCATAACAT
 CCATGAGCGGGGGGACCCCTGGGGGGTGTGGTCTTATGCTGAACCTGAGTATTGTA

 CysSerSerAsnValSerValAlaHisAspGlyAlaGlyLysArgValTyrTyrLeuThr
 6481 CATGCTCTCCAACGTGTGAGTGGCCACGACGGCGCTGGAAAGAGGGTCTACTACCTCA
 GTACGAGGAGGTTGCACAGTCAGCGGGTGTGCGCGACCTTTCTCCAGATGATGGAGT

 ArgAspProThrThrProLeuAlaArgAlaAlaTrpGluThrAlaArgHisThrProVal
 6541 CCGTGACCCTACAACCCCCCTCGCGAGAGCTGCGTGGGAGACAGCAAGACACACTCCAG
 GGGCACTGGGATGTTGGGGGGAGCGCTCTCGACGACCTCTGTGCTTCTGTGTGAGGTC

 AsnSerTrpLeuGlyAsnIleIleMetPheAlaProThrLeuTrpAlaArgMetIleLeu
 6601 TCAATTCCTGGCTAGGCAACATAATCATGTTTGGCCCCACACTGTGGGCGAGGATGATAC
 AGTTAAGGACCGATCCGTTGTATTAGTACAAACGGGGGTGTGACACCGCTCCTACTATG

 MetThrHisPhePheSerValLeuIleAlaArgAspGlnLeuGluGlnAlaLeuAspCys
 6661 TGATGACCATTTCTTTAGCGTCTTATAGCCAGGACAGCTTGAACAGGCCCTCGATT
 ACTACTGGGTAAAGAAATCGCAGGAATATCGGTCCCTGGTGAACCTGTGCGGGAGCTAA

 GluIleTyrGlyAlaCysTyrSerIleGluProLeuAspLeuProProIleIleGlnArg
 6721 GCGAGATCTACGGGGCCTGCTACTCCATAGAACCATTGATCTACCTCCAATCATTCAA
 CGCTCTAGATGCCCCGGACGATGAGGTATCTTGGTGAAC TAGATGGAGGTTAGTAAGTTT

 Leu
 6781 GACTC
 CTGAG

FIG. 32-7

FIG. 47-1 COMBINED ORF OF DNAs K9-1 through 15e

GlyCysProGluArgLeuAlaSerCysArgProLeuThrAspPheAspGlnGlyTrpGly
 1 CAGGCTGCCTGAGAGGCTAGCCAGCTGCCGACCCCTTACCGATTTTGACCAGGGCTGGG
 GTCCGACAGGACTCTCCGATCGGTCGACGGCTGGGGAATGGCTAAACTGGTCCCCGACCC

ProIleSerTyrAlaAsnGlySerGlyProAspGlnArgProTyrCysTrpHisTyrPro
 61 GCCCTATCAGTTATGCCAACGGAAGCGCCCCGACCAGCGCCCTACTGCTGGCACTACC
 CGGGATAGTCAATACGGTTGCCTTCGCCGGGGCTGGTCGCGGGGATGACGACCGTGATGG

ProLysProCysGlyIleValProAlaLysSerValCysGlyProValTyrCysPheThr
 121 CCCCCAAACCTTGCGGTATTGTGCCCGGAAGAGTGTGTGTGGTCCGGTATATTGCTTCA
 GGGGTTTTTGAACGCCATAACACGGGCGCTTCTCACACACACCAGGCCATATAACGAAGT

ProSerProValValValGlyThrThrAspArgSerGlyAlaProThrTyrSerTrpGly
 181 CTCCCAGCCCCGTGGTGGTGGGAACGACCGACAGGTGCGGGCGCGCCACCTACAGCTGGG
 GAGGGTCGGGGCACCACCACCCTTGCTGGCTGTCCAGCCCGCGCGGGTGGATGTGACCC

GluAsnAspThrAspValPheValLeuAsnAsnThrArgProProLeuGlyAsnTrpPhe
 241 GTGAAATGATACGGACGTCTTCGTCTTAACAATACCAGGCCACCGTGGGCAATTGGT
 CACTTTTACTATGCTGCAGAAGCAGGAATTGTTATGGTCCGGTGGCGACCCGTTAACCA

GlyCysThrTrpMetAsnSerThrGlyPheThrLysValCysGlyAlaProProCysVal
 301 TCGGTTGTACCTGGATGAACCTCAACTGGATTACCAAAGTGTGCGGAGCGCCTCCTTGTG
 AGCCAACATGGACCTACTTGAGTTGACCTAAGTGGTTTCACACGCCTCGCGGAGGAACAC

IleGlyGlyAlaGlyAsnAsnThrLeuHisCysProThrAspCysPheArgLysHisPro
 361 TCATCGGAGGGGCGGCAACAACCCCTGCACCTGCCCCACTGATTGCTTCCGCAAGCATC
 AGTAGCCTCCCCGCCGTTGTTGTGGGACGTGACGGGGTACTAACGAAGGCGTTCGTAG

AspAlaThrTyrSerArgCysGlySerGlyProTrpIleThrProArgCysLeuValAsp
 421 CGGACGCCACATACTCTCGGTGCGGCTCCGGTCCCTGGATCACACCCAGGTGCCTGGTTCG
 GCCTGCGGTGTATGAGAGCCACGCCGAGGCCAGGACCTAGTGTGGGTCCACGGACACG

TyrProTyrArgLeuTrpHisTyrProCysThrIleAsnTyrThrIlePheLysIleArg
 481 ACTACCCGTATAGGCTTTGGCATTATCCTTGTACCATCAACTACCCATATTTAAATCA
 TGATGGGCATATCCGAAACCGTAATAGGAACATGGTAGTTGATGTGGTATAAATTTAGT

MetTyrValGlyGlyValGluHisArgLeuGluAlaAlaCysAsnTrpThrArgGlyGlu
 541 GGATGTACGTGGGAGGGGTGGAACACAGGCTGGAAGCTGCCTGCAACTGGACGCGGGGCG
 CCTACATGCACCCTCCCCAGCTTGTGTCCGACCTTCGACGGACGTTGACCTGCGCCCCGC

ArgCysAspLeuGluAspArgAspArgSerGluLeuSerProLeuLeuLeuThrThrThr
 601 AACGTTGCGATCTGGAAGACAGGGACAGGTCCGAGCTCAGCCCGTTACTGCTGACCACTA
 TTGCAACGCTAGACCTTCTGTCCCTGTCCAGGCTCGAGTCGGGCAATGACGACTGGTGAT

GlnTrpGlnValLeuProCysSerPheThrThrLeuProAlaLeuSerThrGlyLeuIle
 661 CACAGTGGCAGGTCTCCCGTGTTCCTTACAACCCCTACCAGCCTTGTCCACCGGCCTCA
 GTGTCAACGCTCAGGAGGGCACAAGGAAGTGTGGGATGGTGGGAACAGGTGGCCGGAGT

HisLeuHisGlnAsnIleValAspValGlnTyrLeuTyrGlyValGlySerSerIleAla
 721 TCCACCTCCACCAGAACATTGTGGACGTGCAGTACTTGTACGGGGTGGGGTCAAGCATCG
 AGGTGGAGGTGGTCTTGTAAACCTGCACGTCAATGAACATGCCCCACCCAGTTCGTAGC

SerTrpAlaIleLysTrpGluTyrValValLeuLeuPheLeuLeuAlaAspAlaArg
 781 CGTCCTGGGCCATTAGTGGGAGTACGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT
 GCAGGACCCGGTAATTCACCCCTATGCAGCAAGAGGACAAGGAAGACGAACGCTCTGCGCG

ValCysSerCysLeuTrpMetMetLeuLeuIleSerGlnAlaGluAlaAlaLeuGluAsn
 841 GCGTCTGCTCCTGCTTGTGGATGATGCTACTCATATCCCAAGCGGAGGCGGCTTTGGAGA
 CGCAGACGAGGACGAACACCTACTACGATGAGTATAGGGTTCGCTCCGCCGAAACCTCT

LeuValIleLeuAsnAlaAlaSerLeuAlaGlyThrHisGlyLeuValSerPheLeuVal
 901 ACCTCGTAATACTTAATGCAGCATCCCTGGCCGGGACGCACGGTCTTGTATCCTTCCTCG
 TGGAGCATTATGAATTACGTCGTAGGGACCGGCCCTGCGTGCCAGAACATAGGAAGGAGC

- 961 PhePheCysPheAlaTrpTyrLeuLysGlyLysTrpValProGlyAlaValTyrThrPhe
TGTTCTTCTGCTTTGTCATGGTATTTGAAGGGTAAGTGGGTGCCCCGAGCGGTCTACACCT
ACAAGAAGACGAAACGTACCATAAACTTCCCATTCACCCACGGGCCCTGCCAGATGTGGA
- 1021 TyrGlyMetTrpProLeuLeuLeuLeuLeuAlaLeuProGlnArgAlaTyrAlaLeu
TCTACGGGATGTGGCCTCTCCTGCTCCTGTTGGCGTTGCCCCAGCGGGCGTACGGC
AGATGCCCTACACCGGAGAGGAGGACGAGGACAACCGCAACGGGGTGCCTCCATGCGCG
- 1081 AspThrGluValAlaAlaSerCysGlyGlyValValLeuValGlyLeuMetAlaLeuThr
TGGACACGGAGGTGGCCGCGTGTGTGGCGGTGTTGTTCTCGTCCGGTTGATGGCGCTGA
ACCTGTGCCTCCACCGGCGCAGCACACCGCCACAACAAGAGCAGCCAACTACCGCGACT
- 1141 LeuSerProTyrTyrLysArgTyrIleSerTrpCysLeuTrpTrpLeuGlnTyrPheLeu
CTCTGTCACCATATTACAAGCGCTATATCAGCTGGTGCTGTGTGGTGGCTTCAGTATTTTC
GAGACAGTGGTATAATGTTCCGATATAGTCGACCACGAACACCACOGAAGTCATAAAAG
- 1201 ThrArgValGluAlaGlnLeuHisValTrpIleProProLeuAsnValArgGlyGlyArg
TGACCAGAGTGAAGCGCAACTGCACGTGTGGATTCCCCCCTCAACGTCCGAGGGGGC
ACTGGTCTACCTTCGGTTGACGTGCACACCTAAGGGGGGAGTTGCAGGCTCCCCCG
- 1261 AspAlaValIleLeuLeuMetCysAlaValHisProThrLeuValPheAspIleThrLys
GCGACGCGTCATCTTACTCATGTGTGTGTACACCGACTCTGGTATTTGACATCACCA
CGTGTGCGCAGTAGAATGAGTACACACGACATGTGGGCTGAGACCATAAACTGTAGTGGT
- 1321 LeuLeuLeuAlaValPheGlyProLeuTrpIleLeuGlnAlaSerLeuLeuLysValPro
AATTGCTGTGGCCGTCTTCGGACCCCTTTGGATTCTTCAAGCCAGTTTGCTTAAAGTAC
TTAACGACGACCGGCAGAACCTGGGGAACCTAAGAAGTTCGGTCAAACGAATTTTCATG
- 1381 TyrPheValArgValGlnGlyLeuLeuArgPheCysAlaLeuAlaArgLysMetIleGly
CCTACTTTGTGGCGCTCCAAGGCCTTCTCCGTTCTGCGCGTTAGCGCGGAAGATGATCG
GGATGAAACACGCGCAGGTTCCGGAAGAGGCCAAGACGCGCAATCGCGCCTTCTACTAGC
- 1441 GlyHisTyrValGlnMetValIleIleLysLeuGlyAlaLeuThrGlyThrTyrValTyr
GAGGCCATTACGTGCAAATGGTCATCATTAAAGTTAGGGGCGCTTACTGGCACCTATGTTT
CTCCGGTAAATGCACGTTTACCAGTAGTAATTCAATCCCGGAATGACCGTGGATACAA
- 1501 AsnHisLeuThrProLeuArgAspTrpAlaHisAsnGlyLeuArgAspLeuAlaValAla
ATAACCATCTCACTCCTCTTCGGGACTGGGCGCACACCGCTTGCAGATCTGGCCGTGG
TATTGGTAGAGTGAGGAGAAGCCCTGACCCGCGTGTGCGGAACGCTCTAGACCGGCACC
- 1561 ValGluProValValPheSerGlnMetGluThrLysLeuIleThrTrpGlyAlaAspThr
CTGTAGAGCCAGTCTCTCTCCCAAATGGAGACCAAGCTCATCACGTGGGGGGCAGATA
GACATCTCGGTACGAGAGAGGGTTTACCTCTGGTTCGAGTAGTGACCCCCCGTCTAT
- 1621 AlaAlaCysGlyAspIleIleAsnGlyLeuProValSerAlaArgArgGlyArgGluIle
CCGCGCGTGGCGTGACATCATCAACGGCTTGCCGTGTTCCGCCCCGAGGGGCCGGGAGA
GGCGGCGCACGCCACTGTAGTAGTTGCCGAACGGACAAAGGGGGCGTCCCCGGCCCTCT
- 1681 LeuLeuGlyProAlaAspGlyMetValSerLysGlyTrpArgLeuLeuAlaProIleThr
TACTGCTGGGGCCAGCCGATGGAATGGTCTCCAAGGGTGGAGGTGCTGGCGCCCATCA
ATGACGAGCCCGGTCCGCTACCTTACCAGAGGTCCCCACCTCCAACGACCGCGGGTAGT
- 1741 AlaTyrAlaGlnGlnThrArgGlyLeuLeuGlyCysIleIleThrSerLeuThrGlyArg
CGGCGTACGCCAGCAGACAAAGGGGCTCTAGGGTGATAATCACCAGCCTAACTGGCC
GCGCATGCGGGTCTGTTTCCCGGAGGATCCACGTATTAGTGGTGGATTGACCGG
- 1801 AspLysAsnGlnValGluGlyGluValGlnIleValSerThrAlaAlaGlnThrPheLeu
GGGACAAAAACCAAGTGGAGGTGAGGTCCAGATTGTGTAAGTCTGCCAAACCTTCC
CCCTGTTTTTGGTTACCTCCCACTCCAGGTCTAACACAGTTGACGACGGGTTTGGAAAG
- 1861 AlaThrCysIleAsnGlyValCysTrpThrValTyrHisGlyAlaGlyThrArgThrIle
TGGCAACGTGCATCAATGGGGTGTGCTGGACTGTCTACCAAGGGGCGGAACGAGGACCA
ACCGTTGCACGTAGTTACCCACACGACCTGACAGATGGTCCCCGGCCTTGCTCCTGGT
- AlaSerProLysGlyProValIleGlnMetTyrThrAsnValAspGlnAspLeuValGly

FIG. 47-2

- 1921 TCGCGTCACCCAAGGGTCCTGTATCCAGATGTATACCAATGTAGACCAAGACCTTGTGG
AGCGCAGTGGGTTCACAGGACAGTAGGTCTACATATGGTTACATCTGGTTCTGGAACACC
- 1981 TrpProAlaProGlnGlySerArgSerLeuThrProCysThrCysGlySerSerAspLeu
GCTGCCCCGCTCCGCAAGGTAGCCGCTCATTGACACCCTGCACCTGCGGCTCCTCGGACC
CGACCGGGCGAGGCGTTCATCGGCGAGTAACTGTGGGACGTGAACGCCGAGGAGCCTGG
- 2041 TyrLeuValThrArgHisAlaAspValIleProValArgArgArgGlyAspSerArgGly
TTTACCTGGTCACGAGGCACGCCGATGTCTATTCCTGCGCCGCGGGGTGATAGCAGGG
AAATGGACCACTGCTCCGTGCGGCTACAGTAAGGGCACGCGGCCGCCCTACTATCGTCCC
- 2101 SerLeuLeuSerProArgProIleSerTyrLeuLysGlySerSerGlyGlyProLeuLeu
GCAGCCTGTGTGCGCCCCGGGCCATTTCTTACTTGAAAGGCTCCTCGGGGGGTCCGCTGT
CGTCGGACGACAGCGGGGCCGGGTAAAGGATGAACCTTCCGAGGAGCCCCCAGGCGACA
- 2161 CysProAlaGlyHisAlaValGlyIlePheArgAlaAlaValCysThrArgGlyValAla
TGTGCCCCGCGGGGCACCGCTGGGCATATTTAGGGCCGCGGTGTGCACCCGTGGAGTGG
ACACGGGGCGCCCCGTGCGGCACCCGTATAAATCCCGCGCCACACGTGGGCACCTCACC
- 2221 LysAlaValAspPheIleProValGluAsnLeuGluThrThrMetArgSerProValPhe
CTAAGGCGGTGGACTTTATCCCTGTGGAGAACCTAGAGACAACCATGAGGTCCCCGGTGT
GATTCGCCACCTGAAATAGGGACACCTCTTGATCTCTGTTGGTACTCCAGGGGCCACA
- 2281 ThrAspAsnSerSerProProValValProGlnSerPheGlnValAlaHisLeuHisAla
TCACGGATAACTCCTCTCCACCACTAGTGCCCCAGAGCTTCCAGGTGGCTCACCTCCATG
AGTGCTATTGAGGAGAGGTGGTCTACACGGGTCTCGAAGGTCCACCGAGTGGAGGTAC
- 2341 ProThrGlySerGlyLysSerThrLysValProAlaAlaTyrAlaAlaGlnGlyTyrLys
CTCCACAGGCAGCGGCAAAAGCACCAAGGTCCCGGCTGCATATGCAGCTCAGGGCTATA
GAGGTGTCCGTGCGCGTTTTCGTGGTTCAGGGCCGACGTATACGTGAGTCCCGATAT
- 2401 ValLeuValLeuAsnProSerValAlaAlaThrLeuGlyPheGlyAlaTyrMetSerLys
AGGTGCTAGTACTCAACCCTCTGTTGTGCAACACTGGGCTTGGTGCTTACATGTCCA
TCCACGATCATGAGTTGGGGAGACAACGACGTTGTGACCCGAAACCAATGTACAGGT
- 2461 AlaHisGlyIleAspProAsnIleArgThrGlyValArgThrIleThrThrGlySerPro
AGGCTCATGGGATCGATCTTAACATCAGGACCGGGGTGAGAACAATTACCACTGGCAGCC
TCCGAGTACCCTAGCTAGGATTGTAGTCTGGCCCCACTCTTGTTAATGGTGACCGTCGG
- 2521 IleThrTyrSerThrTyrGlyLysPheLeuAlaAspGlyGlyCysSerGlyGlyAlaTyr
CCATCAGTACTCCACTACGGCAAGTTCCTTGCGACGGCGGGTGCTCGGGGGGCGCTT
GGTAGTGCATGAGGTGGATGCCGTTCAAGGAACGGCTGCGCCCCAGAGCCCCCGCGAA
- 2581 AspIleIleIleCysAspGluCysHisSerThrAspAlaThrSerIleLeuGlyIleGly
ATGACATAATAATTTGTGACGAGTGCCACTCCAAGGATGCCACATCCATCTTGGGCATCG
TACTGTATTATTAAACACTGCTCACGGTGAGGTGCCTACGGTGTAGGTAGAACCCTAGC
- 2641 ThrValLeuAspGlnAlaGluThrAlaGlyAlaArgLeuValValLeuAlaThrAlaThr
GCACTGTCTTGACCAAGCAGAGACTGCGGGGGCGAGACTGGTTGTGCTCGCCACCGCCA
CGTGACAGGAACCTGGTTCGTCTGACGCCCCGCTCTGACCAACACGAGCGGTGGCGGT
- 2701 ProProGlySerValThrValProHisProAsnIleGluGluValAlaLeuSerThrThr
CCCCTCGGGCTCCGTCACCTGTGCCCCATCCCAACATCGAGGAGGTGCTCTGTCCACCA
GGGGAGGCCCGAGGCAGTGACACGGGGTAGGGTTGTAGCTCCTCCAACGAGACAGGTGGT
- 2761 GlyGluIleProPheTyrGlyLysAlaIleProLeuGluValIleLysGlyGlyArgHis
COGGAGAGATCCCTTTTTACGGCAAGGCTATCCCTCGAAGTAATCAAGGGGGGGAGAC
GGCCTCTCTAGGGAATAATGCCGTTCCGATAGGGGGAGCTTCATTAGTCCCCCCTCTG
- 2821 LeuIlePheCysHisSerLysLysLysCysAspGluLeuAlaAlaLysLeuValAlaLeu
ATCTCATCTTCTGTCAATCAAAGAAGAAGTGCAGCAACTCGCCGCAAAGCTGGTCCGAT
TAGAGTAGAAGACAGTAAGTTTCTTCTTACGCTGCTTGAGCGGCGTTTCGACCAGCGTA
- 2881 GlyIleAsnAlaValAlaTyrTyrArgGlyLeuAspValSerValIleProThrSerGly
TGGGCATCAATGCCGTGGCCTACTACCGCGGTCTTGACGTGTCCGTATCCCGACCAGCG
ACCGTAGTTACGGCACCGGATGATGGCGCCAGAACTGCACAGGCAGTAGGGCTGGTCCG

FIG. 47-3

- 2941 AspValValValValAlaThrAspAlaLeuMetThrGlyTyrThrGlyAspPheAspSer
GCGATGTTGTCGTCGTGGCAACCGATGCCCTCATGACCGGTATACCGGCGACTTCGACT
GGCTACAACAGCAGCACCGTTGGCTACGGGAGTACTGGCCGATATGGCCGCTGAAGCTGA
- 3001 ValIleAspCysAsnThrCysValThrGlnThrValAspPheSerLeuAspProThrPhe
CGGTGATAGACTGCAATACGTGTGTACCCAGACAGTCGATTTCAGCCTTGACCCTACCT
GCCACTATCTGACGTTATGCACACAGTGGGTCTGTCTAGCTAAAGTCGGAACCTGGGATGGA
- 3061 ThrIleGluThrIleThrLeuProGlnAspAlaValSerArgThrGlnArgArgGlyArg
TCACCATTGAGACAATCAGCTCCCCAGGATGCTGTCTCCCGCACTCAACGTCCGGGGCA
AGTGGTAACTCTGTTAGTGGAGGGGGTCTACGACAGAGGGCGTGAGTTGCAGCCCCGT
- 3121 ThrGlyArgGlyLysProGlyIleTyrArgPheValAlaProGlyGluArgProSerGly
GGACTGGCAGGGGAAGCCAGGCATCTACAGATTGTGTGGCAGCGGGGAGCGCCCCCTCCG
CCTGACCGTCCCCCTTCGGTCCGTAGATGTCTAAACACCGTGGCCCCCTCGCGGGGAGGC
- 3181 MetPheAspSerSerValLeuCysGluCysTyrAspAlaGlyCysAlaTrpTyrGluLeu
GCATGTTGACTCGTCCGTCTCTGTGAGTGCTATGACGCAGGCTGTCTTGGTATGAGC
CGTACAAGCTGAGCAGGCAGGAGACACTACGATACTGCGTCCGACACGAACCATACTCG
- 3241 ThrProAlaGluThrThrValArgLeuArgAlaTyrMetAsnThrProGlyLeuProVal
TCACGCCCCGCGAGACTACAGTTAGGCTACGAGCGTACATGAACCCCCGGGGCTTCCCG
AGTGGGGCGGCTCTGATGTCAATCCGATGCTCGCATGTACTTGTGGGGCCCCGAAGGGC
- 3301 CysGlnAspHisLeuGluPheTrpGluGlyValPheThrGlyLeuThrHisIleAspAla
TGTGCCAGGACCATCTTGAATTTGGGAGGGCGTCTTTACAGGCCTCACTCATATAGATG
ACACGCTCTGGTAGAACTTAAACCTCCCGCAGAAATGTCCGGAGTGAGTATATCTAC
- 3361 HisPheLeuSerGlnThrLysGlnSerGlyGluAsnLeuProTyrLeuValAlaTyrGln
CCCACCTTCTATCCAGACAAAGCAGAGTGGGGAGAACCTTCCTTACCTGGTAGCGTACC
GGGTGAAAGATAGGGTCTGTTTCGTCTCACCCCTCTTGAAGGAATGGACCATCGCATGG
- 3421 AlaThrValCysAlaArgAlaGlnAlaProProProSerTrpAspGlnMetTrpLysCys
AAGCCACCGTGTGCGCTAGGGCTCAAGCCCTCCCCCATCGTGGGACCAGATGTGGAAGT
TTCGGTGGCACACGCGATCCCGAGTTCGGGGAGGGGTAGCACCCCTGGTCTACACCTTCA
- 3481 LeuIleArgLeuLysProThrLeuHisGlyProThrProLeuLeuTyrArgLeuGlyAla
GTTTGATTCGCTCAAGCCACCCCTCCATGGGCCAACCCCTGCTATACAGACTGGGGC
CAAACCTAAGCGGAGTTCGGGTGGGAGGTACCCGGTTGTGGGGACGATATGTCTGACCCGC
- 3541 ValGlnAsnGluIleThrLeuThrHisProValThrLysTyrIleMetThrCysMetSer
CTGTTCAAGATGAAATCACCTGACGCACCCAGTCACCAAATACATCATGACATGCATGT
GACAAGTCTTACTTTAGTGGGACTGCGTGGGTCAGTGGTTTATGTAGTACTGTACGTACA
- 3601 AlaAspLeuGluValValThrSerThrTrpValLeuValGlyGlyValLeuAlaAlaLeu
CGGCCGACCTGGAGGTGCTACGAGCACCTGGGTGCTCGTTGGCGGCGTCTGGCTGCTT
GCCGGCTGGACCTCCAGCAGTGCTCGTGACCCACGAGCAACCGCCGAGGACCGACGAA
- 3661 AlaAlaTyrCysLeuSerThrGlyCysValValIleValGlyArgValValLeuSerGly
TGGCCGCTATTGCCTGTCAACAGGCTGCGTGGTCATAGTGGGCAGGGTCTGCTTGTCCG
ACCGGCGCATAACGGACAGTTGTCCGACGCACAGTATACCCGTCCAGCAGAACAGGC
- 3721 LysProAlaIleIleProAspArgGluValLeuTyrArgGluPheAspGluMetGluGlu
GGAAGCCGGCAATCATACCTGACAGGGAAGTCTCTACCGAGAGTTCGATGAGATGGAAG
CCTTCGGCCGTTAGTATGGACTGTCCCTCAGGAGATGGCTCTCAAGCTACTCTACCTTC
- 3781 CysSerGlnHisLeuProTyrIleGluGlnGlyMetMetLeuAlaGluGlnPheLysGln
AGTGCTCTCAGCACTTACCGTACATCGAGCAAGGGATGATGCTCGCCGAGCAGTTCAAGC
TCACGAGAGTCTGTAATGGCATGTAGCTCGTTCCTACTACGAGCGGCTCGTCAAGTTCCG
- 3841 LysAlaLeuGlyLeuLeuGlnThrAlaSerArgGlnAlaGluValIleAlaProAlaVal
AGAAGGCCCTCGGCCTCTGCGAGACCGGCTCCCGTCAGGCAGAGGTATCGCCCCCTGCTG
TCTTCCGGGAGCCGGAGGACGTCTGGCGCAGGGCAGTCTGCTCCAATAGCGGGGACGAC
- GlnThrAsnTrpGlnLysLeuGluThrPheTrpAlaLysHisMetTrpAsnPheIleSer

FIG. 47-4

- 3901 TCCAGACCAACTGGCAAACTCGAGACCTTCTGGGCGAAGCATATGTGGAACCTTCATCA
AGGCTCGGTGACCGTTTTTGAGCTCTGGAAGACCGCTTCGTATACACCTTGAAGTAGT
- GlyIleGlnTyrLeuAlaGlyLeuSerThrLeuProGlyAsnProAlaIleAlaSerLeu
3961 GTGGGATACAATACTTGGCGGGCTGTCAACGCTGCCTGGTAACCCGCCATTGCTTCAT
CACCTATGTTATGAACCGCCGAACAGTTGCGACGGACCATTTGGGGCGTAACGAAGTA
- MetAlaPheThrAlaAlaValThrSerProLeuThrThrSerGlnThrLeuLeuPheAsn
4021 TGATGGCTTTTACAGCTGCTGTACCAGCCACTAACCCTAGCCAAACCTCTCTCA
ACTACCGAAAATGTCGACGACAGTGGTGGGTGATTGGTGTATCGGTTTGGGAGGAGAAGT
- IleLeuGlyGlyTrpValAlaAlaGlnLeuAlaAlaProGlyAlaAlaThrAlaPheVal
4081 ACATATTGGGGGGTGGGTGGCTGCCCAGCTCGCCGCCCCGGTGGCGCTACTGCCTTTG
TGATAACCCCCCACCACCGACGGGTGAGCGCGGGGGCCACGGCGATGACGGAAC
- GlyAlaGlyLeuAlaGlyAlaAlaIleGlySerValGlyLeuGlyLysValLeuIleAsp
4141 TGGGCGCTGGCTTAGCTGGCGCCGCGCATCGGCAGTGTGGACTGGGGAAGGTCTTCATAG
ACCCGCGACCGAATCGACCGCGGGTAGCCGTCAACCTGACCCCTTCCAGGAGTATC
- IleLeuAlaGlyTyrGlyAlaGlyValAlaGlyAlaLeuValAlaPheLysIleMetSer
4201 ACATCCTTGCAGGGTATGCGCGGGCGTGGCGGGAGCTCTTGTGGCATTCAAGATCATGA
TGTAGGAACGTCCCATACCGCGCCCGACCGCCCTCGAGAACACCGTAAGTTCTAGTACT
- GlyGluValProSerThrGluAspLeuValAsnLeuLeuProAlaIleLeuSerProGly
4261 GCGGTGAGGTCCCCTCCACGGAGGACCTGGTCAATCTACTGCCCGCCATCTCTCGCCCG
CGCCACTCCAGGGGAGGTGCTCTGGACAGTTAGATGACGGGCGGTAGGAGAGCGGGC
- AlaLeuValValGlyValValCysAlaAlaIleLeuArgArgHisValGlyProGlyGlu
4321 GAGCCCTCGTAGTCGGCTGGTCTGTGCAGCAATACTGCGCCGGCACGTTGGCCCGGGCG
CTCGGGAGCATCAGCCGCACCAGACACGTCGTTATGACGCGCGCGTCAACCGGGCCCGC
- GlyAlaValGlnTrpMetAsnArgLeuIleAlaPheAlaSerArgGlyAsnHisValSer
4381 AGGGGGCAGTGCAGTGGATGAACCGGCTGATAGCCTTCGCCTCCCGGGGAACCATGTTT
TCCCCCGTCACGTCACCTACTTGGCCGACTATCGGAAGCGGAGGGCCCCCTTGGTACAAA
- ProThrHisTyrValProGluSerAspAlaAlaAlaArgValThrAlaIleLeuSerSer
4441 CCCCCACGCACTACGTGCCGGAGAGCGATGCAGCTGCCCGCGTCACCTGCCTACTCAGCA
GGGGGTGCGTGATGCACGGCCTCTCGCTACGTGCAGGGGCGCAGTGACGGTATGAGTCGT
- LeuThrValThrGlnLeuLeuArgArgLeuHisGlnTrpIleSerSerGluCysThrThr
4501 GCCTCACTGTAAACCCAGCTCCTGAGGCGACTGCACCACTGGATAAGCTCGGAGTGATACCA
CGGAGTGACATTGGGTGCGAGGACTCGCTGACGTGGTACCTATTGAGCCTCACATGGT
- ProCysSerGlySerTrpLeuArgAspIleTrpAspTrpIleCysGluValLeuSerAsp
4561 CTCCATGCTCCGGTTCCCTGGCTAAGGGACATCTGGGACTGGATATGCGAGGTGTTGAGCG
GAGGTACGAGGCCAAGGACCGATTCCCTGTAGACCCCTGACCTATACGCTCCACAACCTCGC
- PheLysThrTrpLeuLysAlaLysLeuMetProGlnLeuProGlyIleProPheValSer
4621 ACTTTAAGACCTGGCTAAAAGCTAAGCTCATGCCACAGCTGCCTGGGATCCCCCTTTGTGT
TGAAATTCTGGACCGATTTCGATTGAGTACGGTGTGACGGACCCTAGGGGAAACACA
- CysGlnArgGlyTyrLysGlyValTrpArgValAspGlyIleMetHisThrArgCysHis
4681 CCTGCCAGCGCGGTATAAGGGGGTCTGGCGAGTGGACGGCATCATGCACACTCGCTGCC
GGACGGTTCGCGCCATATTCCCCCAGACCGCTCACCTGCCGTAGTACGTGTGAGCGACGG
- CysGlyAlaGluIleThrGlyHisValLysAsnGlyThrMetArgIleValGlyProArg
4741 ACTGTGGAGCTGAGATCACTGGACATGTCAAAAACGGGACGATGAGGATCGTCCGTCTTA
TGACACCTCGACTCTAGTGACCTGTACAGTTTTTGGCCCTGCTACTCTAGCAGCCAGGAT
- ThrCysArgAsnMetTrpSerGlyThrPheProIleAsnAlaTyrThrThrGlyProCys
4801 GGACCTGCAGGAACATGTGGAGTGGGACCTTCCCCATTATGCCTACACCACGGGCCCCCT
CCTGGACGTCTTGTACACCTACCCCTGGAAGGGGTAATTACGGATGTGGTGGCCCGGGGA
- ThrProLeuProAlaProAsnTyrThrPheAlaLeuTrpArgValSerAlaGluGluTyr
4861 GTACCCCCCTTCCCTGCGCCGAACCTACAGTTTCGCGCTATGGAGGGTGTCTGCAGAGGAAT
CATGGGGGAAGGACGCGGCTTGATGTCAAGCGGATACCTCCACAGACGTCTCCTTA

FIG. 47-5

4921 ValGluIleArgGlnValGlyAspPheHisTyrValThrGlyMetThrThrAspAsnLeu
 ATGTGGAGATAAGGCAGGTGGGGGACTTCCACTACGTGACGGGTATGACTACTGACAATC
 TACACCTCTATTCCGTCCACCCCTGAAGGTGATGCAC TGCCATACTGATGACTGTTAG

4981 LysCysProCysGlnValProSerProGluPhePheThrGluLeuAspGlyValArgLeu
 TCAATGCCCGTGCCAGGTCCCATCGCCCGAATTTTTCACAGAATTGGACGGGGTGCGCC
 AGTTTACGGGCACGGTCCAGGTAGCGGGCTTAAAAAGTGCTTAACCTGCCCCACGCGG

5041 HisArgPheAlaProProCysLysProLeuLeuArgGluGluValSerPheArgValGly
 TACATAGGTTTGC GCCCCCCCTGCAAGCCCTTGCTGCGGGAGGAGGTATCATTGAGAGTAG
 ATGTATCCAAACGCGGGGGGACGTTCGGGAACGACGCCCTCTCCATAGTAAGTCTCATC

5101 LeuHisGluTyrProValGlySerGlnLeuProCysGluProGluProAspValAlaVal
 GACTCCACGAATACCCGGTAGGGTCGCAATTACCTTGCGAGCCCGAACCGGACGTGGCCG
 CTGAGGTGCTTATGGGCCATCCAGCGTTAATGGAACGCTCGGGCTTGCCCTGCACCGGC

5161 LeuThrSerMetLeuThrAspProSerHisIleThrAlaGluAlaAlaGlyArgArgLeu
 TGTGTAGCTCCATGCTCACTGATCCCTCCCATATAACAGCAGAGGCGGGCGGGCGAAGGT
 ACAACTGCAGGTACGAGTACTAGGGAGGGTATATTGTCGTCTCCGCCGGCCCGCTTCCA

5221 AlaArgGlySerProProSerValAlaSerSerSerAlaSerGlnLeuSerAlaProSer
 TGGCGAGGGGATCACCCCTCTGTGGCCAGCTCCTCGGCTAGCCAGCTATCCGCTCCAT
 ACCGCTCCCTAGTGGGGGAGACACCGGTGAGGAGCCGATCGGTGATAGGCGAGGTA

5281 LeuLysAlaThrCysThrAlaAsnHisAspSerProAspAlaGluLeuIleGluAlaAsn
 CTCTCAAGCAACTTGCACCGCTAACCATGACTCCCTGATGCTGAGCTCATAGAGGCCA
 GAGAGTTCGGTTGAACGTGGCGATTGGTACTGAGGGGACTACGACTCGAGTATCTCCGGT

5341 LeuLeuTrpArgGlnGluMetGlyGlyAsnIleThrArgValGluSerGluAsnLysVal
 ACCTCTATGGAGGCAGGAGATGGGCGGCAACATCACCAGGGTTGAGTCAGAAAACAAG
 TGGAGGATACCTCCGTCTCTACCCGCCGTGTAGTGGTCCCAACTCAGTCTTTTGTTC

5401 ValIleLeuAspSerPheAspProLeuValAlaGluGluAspGluArgGluIleSerVal
 TGGTGATTCTGGACTCCTTCGATCCGCTTGTTGGCGGAGGAGACGAGCGGGAGATCTCCG
 ACCACTAAGACCTGAGGAAGCTAGGCGAACACCGCCTCTCTGCTCGCCCTCTAGAGGC

5461 ProAlaGluIleLeuArgLysSerArgArgPheAlaGlnAlaLeuProValTrpAlaArg
 TACCCGACAGAAATCTGCGGAAGTCTCGGAGATTGCGCCAGGCCCTGCCGTTTGGGGCGC
 ATGGGCGTCTTTAGGACGCCCTCAGAGCCTTAAGCGGGTCCGGGACGGGCAACCCCGC

5521 ProAspTyrAsnProProLeuValGluThrTrpLysLysProAspTyrGluProProVal
 GGCCGGACTATAACCCCGCTAGTGGAGACGTGGAAAAAGCCGACTACGAACCACCTG
 CCGCCTGATATTGGGGGGGATCACCTCTGCACCTTTTTCGGGCTGATGCTTGGTGGAC

5581 ValHisGlyCysProLeuProProProLysSerProProValProProProArgLysLys
 TGGTCCATGGCTGTCCGCTTCCACCTCAAAGTCCCTCTGTGCTCCGCTCGGAAGA
 ACCAGGTACCGACAGGCGAAGGTGGAGGTTTCAGGGGAGGACACGGAGGCGGAGCCTTCT

5641 ArgThrValValLeuThrGluSerThrLeuSerThrAlaLeuAlaGluLeuAlaThrArg
 AGCGGACGGTGGTCTCTCACTGAATCAACCTATCTACTGCCTTGGCCGAGCTCGCCACCA
 TCGCTGCCACAGGAGTACTTAGTGGGATAGATGACGGAACCGGCTCGAGCGGTGGT

5701 SerPheGlySerSerSerThrSerGlyIleThrGlyAspAsnThrThrThrSerSerGlu
 GAAGCTTTGGCAGCTCCTCAACTTCCGGCATTACGGGCGACAATACGACAACATCCTCTG
 CTTGAAACCGTCGAGGAGTTGAAGGCCGTAATGCCGCTGTTATGCTGTTGTAGGAGAC

5761 ProAlaProSerGlyCysProProAspSerAspAlaGluSerTyrSerSerMetProPro
 AGCCCGCCCTTCTGGCTGCCCCCGACTCCGACGCTGAGTCTTATCTCCATGCCCC
 TCGGGCGGGGAAGACCGACGGGGGGCTGAGGCTGCGACTCAGGATAAGGAGGTACGGG

5821 LeuGluGlyGluProGlyAspProAspLeuSerAspGlySerTrpSerThrValSerSer
 CCCTGAGGGGGAGCCTGGGATCCGGATCTTAGCGACGGGTCATGGTCAACGGTCAGTA
 GGGACCTCCCCCTCGGACCCCTAGGCTAGAATCGCTGCCAGTACCAGTTGCCAGTCAT

GluAlaAsnAlaGluAspValValCysCysSerMetSerTyrSerTrpThrGlyAlaLeu

FIG. 47-6

5881 GTGAGGCCAACGCGGAGGATGTCGTGTGCTGCTCAATGTCTTACTCTTGGACAGGCGCAC
 CACTCCGGTTGCGCCTCTACAGCACACGACGAGTTACAGAATGAGAACCTGTCCGCGTG

 ValThrProCysAlaAlaGluGluGlnLysLeuProIleAsnAlaLeuSerAsnSerLeu
 5941 TCGTACCCCCGTGCGCCGCGGAAGAACAGAACTGCCCATCAATGCACTAAGCAACTCGT
 AGCAGTGGGGCACGCGGCGCCTTCTTGTCTTTGACGGGTAGTTACGTGATTCTGTTGAGCA

 LeuArgHisHisAsnLeuValTyrSerThrThrSerArgSerAlaCysGlnArgGlnLys
 6001 TGCTACGTACCCACAATTGGTGTATTCCACCACCTCACGCAGTGTGTCGCAAAGGCAGA
 ACATGTCAGTGGTGTAAACCACATAAGGTGGTGGAGTGCCTCACGAACGGTTTCCGTCT

 LysValThrPheAspArgLeuGlnValLeuAspSerHisTyrGlnAspValLeuLysGlu
 6061 AGAAAGTCACATTGACAGACTGCAAGTTCGACAGCCATTACCAGGACGTACTCAAGG
 TCTTTCAGTGTAACCTGTCTGACGTTCAAGACCTGTCGGTAATGGTCTGTCATGAGTTCC

 ValLysAlaAlaAlaSerLysValLysAlaAsnLeuLeuSerValGluGluAlaCysSer
 6121 AGGTTAAAGCAGCGCGCTCAAAAGTGAAGGCTAACTTGCTATCCGTAGAGGAAGCTTGCA
 TCCAATTTCTGTCGCCGCGAGTTTCACTTCCGATTGAACGATAGGCATCTCCTTCGAACGT

 LeuThrProProHisSerAlaLysSerLysPheGlyTyrGlyAlaLysAspValArgCys
 6181 GCCTGACGCCCCACACTCAGCCAAATCCAAGTTTGGTTATGGGGCAAAGACGTCGGTT
 CGGACTGCGGGGGTGTGAGTCGGTTTAGGTTCAAACCAATACCCGTTTCTGTCAGGCAA

 HisAlaArgLysAlaValThrHisIleAsnSerValTrpLysAspLeuLeuGluAspAsn
 6241 GCCATGCCAGAAAGGCCGTAAACCCACATCAACTCCGTGTGGAAAGACCTTCTGGAAGACA
 CGGTACGGTCTTTCCGGCATTGGGTGTAGTTGAGGCACACCTTCTGGAAGACCTTCTGT

 ValThrProIleAspThrThrIleMetAlaLysAsnGluValPheCysValGlnProGlu
 6301 ATGTAAACCAATAGACACTACCATCATGGCTAAGAACGAGGTTTTCTGCGTTCAGCCTG
 TACATTGTGGTTATCTGTGATGGTAGTACCGATTCTTGCTCCAAAGACGCAAGTCGGAC

 LysGlyGlyArgLysProAlaArgLeuIleValPheProAspLeuGlyValArgValCys
 6361 AGAAGGGGGGTGCTAAGCCAGCTCGTCTCATCGTGTTCCTCCGATCTGGGCGTGCGCGTGT
 TCTTCCCCCAGCATTCCGTCGAGCAGAGTAGCACAAGGGGCTAGACCCGCACGCGCACA

 GluLysMetAlaLeuTyrAspValValThrLysLeuProLeuAlaValMetGlySerSer
 6421 GCGAAAGATGGCTTTGTACGACGTGGTTACAAAGCTCCCTTGGCCGTGATGGGAAGCT
 CGCTTTTCTACCGAAACATGCTGCACCAATGTTTCGAGGGGAACCGGCACTACCCTTCGA

 TyrGlyPheGlnTyrSerProGlyGlnArgValGluPheLeuValGlnAlaTrpLysSer
 6481 CCTACGGATTCCAATACTCACCAGACAGCGGGTTGAATTCCTCGTGCAAGCGTGGAAGT
 GGATGCCAAGGTTATGAGTGGTCTGTCGCCCACTTAAGGAGCACGTTCCGACCTTCA

 LysLysThrProMetGlyPheSerTyrAspThrArgCysPheAspSerThrValThrGlu
 6541 CCAAGAAACCCCAATGGGGTCTCGTATGATACCCGCTGCTTGACTCCACAGTCACTG
 GGTCTTTTGGGGTTACCCCAAGAGCATACTATGGGCGACGAAACTGAGGTGTCAGTGAC

 SerAspIleArgThrGluGluAlaIleTyrGlnCysCysAspLeuAspProGlnAlaArg
 6601 AGAGCGACATCCGTACGGAGGAGGCAATCTACCAATGTTGTGACCTCGACCCCAAGCCC
 TCTCGCTGTAGGCATGCCTCCTCCGTTAGATGGTTACAACACTGGAGCTGGGGGTTCCGG

 ValAlaIleLysSerLeuThrGluArgLeuTyrValGlyGlyProLeuThrAsnSerArg
 6661 GCGTGGCCATCAAGTCCCTCACCAGAGAGGCTTATGTTGGGGGCCCTCTTACCAATTCAA
 CGCACCGGTAGTTTACGGAGTGGCTCTCCGAAATACAACCCCGGGAGAATGGTTAAGTT

 GlyGluAsnCysGlyTyrArgArgCysArgAlaSerGlyValLeuThrThrSerCysGly
 6721 GGGGGGAGAACTGCGGCTATCGCAGGTGCGCGCGAGCGGCTACTGACAACTAGCTGTG
 CCCCCCTTTGACGCCGATAGCGTCCACGGCGCGCTCGCCGATGACTGTTGATCGACAC

 AsnThrLeuThrCysTyrIleLysAlaArgAlaAlaCysArgAlaAlaGlyLeuGlnAsp
 6781 GTAACACCTCACTTGCTACATCAAGGCCGGGCAGCCTGTGAGCCGCGAGGGCTCCAGG
 CATGTGGGAGTGAACGATGTAGTTCCGGGCCCGTCCGACAGCTCGGCGTCCCGAGGTCC

 CysThrMetLeuValCysGlyAspAspLeuValValIleCysGluSerAlaGlyValGln
 6841 ACTGCACCATGCTCGTGTGTGGCGACGACTAGTCGTTATCTGTGAAAGCGCGGGGGTCC
 TGACGTGGTACGAGCACACACCGCTGCTGAATCAGCAATAGACACTTTCGCGCCCCCAGG

FIG. 47-7

6901 GluAspAlaAlaSerLeuArgAlaPheThrGluAlaMetThrArgTyrSerAlaProPro
AGGAGGACGCGGCGAGCCTGAGAGCCTTCACGGAGGCTATGACCAGGTACTCCGCCCCC
TCCTCCTGCGCGCTCGGACTCTCGGAAGTGCTCCGATACTGGTCCATGAGGCGGGGG

6961 GlyAspProProGlnProGluTyrAspLeuGluLeuIleThrSerCysSerSerAsnVal
CTGGGGACCCCCACAACCAGAATACGACTTGGAGCTCATAACATCATGCTCCTCCAACG
GACCCCTGGGGGGTGTGGTCTTATGCTGAACCTCGAGTATTGTAGTACGAGGAGGTGC

7021 SerValAlaHisAspGlyAlaGlyLysArgValTyrTyrLeuThrArgAspProThrThr
TGTCAGTCGCCCACGACGGCGCTGGAAAGAGGGTCTACTACCTACCCGTGACCCACAA
ACAGTCAGCGGGTGCTGCCGCGACCTTTCTCCAGATGATGGAGTGGGCACTGGGATGTT

7081 ProLeuAlaArgAlaAlaTrpGluThrAlaArgHisThrProValAsnSerTrpLeuGly
CCCCCTCGCGAGAGCTGCGTGGGAGACAGCAAGACACACTCCAGTCAATTCCTGGCTAG
GGGGGAGCGCTCTCGACGCACCCTCTGTCGTTCTGTGTGAGGTCAGTTAAGGACCGATC

7141 AsnIleIleMetPheAlaProThrLeuTrpAlaArgMetIleLeuMetThrHisPhePhe
GCAACATAATCATGTTTGCCCCACACTGTGGGCGAGGATGATACTGATGACCCATTCT
CGTGTATTAGTACAAACGGGGGTGTGACACCCGCTCCTACTATGACTACTGGGTAAAGA

7201 SerValLeuIleAlaArgAspGlnLeuGluGlnAlaLeuAspCysGluIleTyrGlyAla
TTAGCGTCCTTATAGCCAGGGACAGCTTGAACAGGCCCTCGATTGCGAGATCTACGGGG
AATCGCAGGAATATCGGTCCCTGGTGAACCTGTCCGGGAGCTAACGCTCTAGATGCCCC

7261 CysTyrSerIleGluProLeuAspLeuProProIleIleGlnArgLeu
CCTGCTACTCCATAGAACCCTTGATCTACCTCCAATCATTCAAAGACTC
GGACGATGAGGTATCTTGGTGAACCTAGATGGAGGTTAGTAAGTTTCTGAG

FIG. 47- 8